Detecting Recessive Diseases in Inbred Populations

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ABSTRACT OF THE DISCLOSURE

Techniques of using statistical analysis of genetic data to determine likely markers for a recessive genetic disease or trait. One embodiment of these techniques includes the steps of obtaining actual genotype data for one or more affected people with the genetic disease or trait in a population, obtaining estimated genotype data for the population, and analyzing the actual and estimated genotype data to find a region in genomes of the affected people that includes markers exhibiting particular homozygous pairs of alleles more frequently than would occur randomly.

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